#### **Brief Communication**

# Involvement of the Lateral Hypothalamic Peptide Orexin in Morphine Dependence and Withdrawal

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The lateral hypothalamus (LH) is implicated in the behavioral actions of drugs of abuse, but the cellular and molecular basis of this role is unclear. Recent identification of neuropeptides localized in LH neurons has allowed for more specific studies of LH function. The LH-specific peptide orexin (hypocretin) has been shown to be important in arousal and sleep regulation. However, orexin cells of the LH project broadly throughout the brain such that orexin may influence other behaviors as well. In this study, we show that orexin neurons, and not nearby LH neurons expressing melanin-concentrating hormone (MCH), have  $\mu$ -opioid receptors and respond to chronic morphine administration and opiate antagonist-precipitated morphine withdrawal. cAMP response element-mediated transcription is induced in a subset of orexin cells, but not MCH cells, after exposure to chronic morphine or induction of withdrawal. Additionally, c-Fos and the orexin gene itself are induced in orexin cells in the LH during morphine withdrawal. Finally, we show that orexin knock-out mice develop attenuated morphine dependence, as indicated by a less severe antagonist-precipitated withdrawal syndrome. Together, these studies support a role for the orexin system in molecular adaptations to morphine, and demonstrate dramatic differences in molecular responses among different populations of LH neurons.

Key words: μ-opioid receptor; CREB; c-Fos; melanin-concentrating hormone; opiate withdrawal; drug addiction

### Introduction

Lesion and intracranial self-stimulation (ICSS) studies have suggested an important role for the lateral hypothalamus (LH) in feeding, arousal, and reward (Anand and Brobeck, 1951; Olds and Milner, 1954). Compared with other brain regions, ICSS in the LH is the most robust (Gallistel et al., 1981) and can be modulated by opiates and several other drugs of abuse or their antagonists (Adams et al., 1972; Goodall and Carey, 1975; Bozarth et al., 1980). Opiates, such as morphine, are also self-administered directly into the LH (Olds and Williams, 1980; Cazala et al., 1987). The target of these selfstimulation and self-administration studies is a subregion of the LH that is crossed by the medial forebrain bundle (MFB). Besides the passing fibers that connect limbic structures from the basal forebrain with the midbrain, the MFB also contains axons of intrinsic LH neurons (Millhouse, 1969). Moreover, previous studies have shown that ibotenic acid lesions of the LH, which destroy cell bodies but leave the passing MFB fibers intact, decrease ICSS in the LH (Velley et al., 1983). However, little is known about the specific neuronal cell types in the LH that contribute to these behaviors.

Recently, several neuropeptides have been shown to be highly enriched in the LH. Two neuropeptides, orexin (also called hypocretin) and melanin-concentrating hormone (MCH), are expressed in nonoverlapping populations of LH neurons (Peyron et al., 1998). Knock-out and transgenic overexpression studies have shown that MCH is important in feeding (Ludwig et al., 1998; Shimada et al., 1998), whereas orexin is important in arousal and sleep (Chemelli et al., 1999). However, there is evidence that orexin neurons may regulate other behaviors, such as stress responses (Ida et al., 2000) and feeding (Edwards et al., 1999; Haynes et al., 1999). This is consistent with the fact that orexin neurons project to many brain areas. Among these orexin projection regions are several that are implicated in behavioral responses to drugs of abuse, such as the locus ceruleus, nucleus accumbens, and ventral tegmental area (Peyron et al., 1998; Fadel and Deutch, 2002). However, little is known about the function of orexin neurons in influencing responses to drugs of abuse. In this study, we use cAMP response element (CRE) reporter mice, orexin gene reporter mice, and orexin knock-out mice to demonstrate a role for orexin in regulating morphine dependence and withdrawal.

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### **Materials and Methods**

Transgenic mouse lines. We used CRE-LacZ reporter (line 37), orexin knock-out (Impey et al., 1998), and orexin reporter (orexin-τ-LacZ) mice (Chemelli et al., 1999) that had all been backcrossed more than five times to a C57BL/6 background. Animals were bred and maintained under a 12

hr dark/light cycle with food and water available *ad libitum* and were used at 8–12 weeks of age; littermates were used as controls in all experiments.

Behavioral scoring. Homozygous male orexin knock-out mice and wild-type littermates were implanted with a 25 mg morphine base pellet subcutaneously, or received sham surgery (no pellet), under light isofluorane anesthesia. Two days later, mice received naloxone (1 mg/kg, s.c.), and withdrawal symptoms (jumps, wet dog shakes, backwards walking, paw tremor, tremor, diarrhea, ptosis, and weight loss) were monitored for 25 min (Shaw-Lutchman et al., 2002). A global withdrawal score was calculated by multiplying withdrawal signs by a constant and adding the scores for each sign. A separate set of mice was exposed to an open field, and activity and locomotion were recorded for 3 hr before the onset of the dark cycle.

Immunohistochemical studies. Orexin- $\tau$ -LacZ and CRE-LacZ mice were implanted subcutaneously on day 1 with a 25 mg morphine pellet or received sham surgery. An identical procedure was performed on day 3. On day 6, mice received saline or naltrexone (100 mg/kg, s.c.), a dose required for maximal, sustained levels of withdrawal (Shaw-Lutchman et al., 2002). Four hours later (to permit reporter gene expression), animals were injected with an overdose of pentobarbital and perfused transcardially with 1× PBS followed by 4% paraformaldehyde. The brains were postfixed for 12 hr in 4% paraformaldehyde and then cryoprotected in 20% glycerol for 6 hr. Brains were sectioned at 40  $\mu$ m intervals and collected in 1× PBS plus 0.05% sodium azide.

Double-labeling immunohistochemistry was performed as described previously (Shaw-Lutchman et al., 2002) using goat polyclonal anti- $\beta$ -galactosidase ( $\beta$ -gal) antibody (1:5000; Biogenesis, Poole, UK), rabbit polyclonal anti-orexin antibody (1:5000; Ghemicon, Temecula, CA), rabbit polyclonal anti-MCH antibody (1:5000; gift from W. Vale, Salk Institute, La Jolla, CA), goat polyclonal anti-c-Fos antibody (1:500; Santa Cruz Biotechnology, Santa Cruz, CA), goat polyclonal anti-FosB antibody (1:500; Santa Cruz Biotechnology), or rabbit polyclonal anti- $\mu$ -opioid receptor (anti-MOR; 1:4000; ImmunoStar, Hudson, WI). Quantification and localization of  $\beta$ -gal, c-Fos, or FosB expression with orexin or MCH were performed using fluorescent light microscopy by an investigator blinded to treatment conditions.

Triple-labeling immunocytochemistry used mouse anti-orexin (1:500; R&D Systems, Minneapolis, MN), goat anti-c-Fos (1:500) or goat anti- $\beta$ -gal (1:5000), and rabbit anti- $\mu$ -opioid receptor (1:4000). Secondary antibodies were added sequentially: first, the Cy-2 donkey anti-goat (1:200) and Cy-5 donkey anti-mouse antibodies in 1× PBS for 4 hr, followed by biotinylated goat anti-rabbit antibody (1:400) in 1× PBS for 1 hr. An avidin-biotin complex (ABC) reaction (Vectastain Elite kit; Vector Laboratories, Burlingame, CA) was performed for 10 min followed by a 10 min tyramide signal amplification (TSA Plus Fluorescence Systems kit; PerkinElmer Life Sciences, Boston, MA). Each of the four components of the amplification reaction was omitted as a control. Analysis of triple-labeling was performed by confocal microscopy (LSM 510; Zeiss, Oberkochen, Germany) at 630× magnification. Sections were optically sliced in the *z*-plane at 0.4  $\mu$ m intervals.

Analysis of  $\beta$ -gal expression. Perfused brains of orexin- $\tau$ -LacZ mice were cut as 40  $\mu$ m coronal sections, and 5-bromo-4-chloro-3-indolyl- $\beta$ -D-galactopyranoside histochemistry was performed as described previously (Min et al., 1994). For quantification, images of brain sections were captured at 200× magnification, and densitometry was performed on anteroposteriorly matched sections using NIH Image software, as described previously (Min et al., 1996). For every mouse used (six per group), the density of all cell bodies (5–15 per section) from three adjacent, anatomically matched sections was measured and averaged.

Statistical analysis. Results were expressed as mean  $\pm$  SEM. Statistical significance was estimated using the t test or one-way ANOVA followed by the Tukey post hoc test when appropriate. A p value of <0.05 was considered significant.

# Results

# Orexin neurons are regulated by chronic morphine and morphine withdrawal

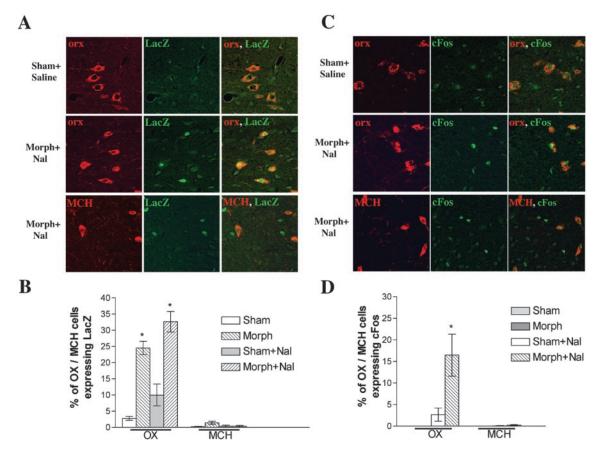
To better understand the effects of morphine on the LH, we evaluated molecular responses to chronic morphine and mor-

phine withdrawal precipitated by the opioid receptor antagonist naltrexone. CRE-LacZ transgenic mice were used as a reporter of CRE-mediated transcription (Impey et al., 1998), which is known to be regulated by morphine in several other brain regions (Shaw-Lutchman et al., 2002). In these mice, the LacZ gene is induced with activation of CRE-binding protein (CREB) or a CREB family protein (Montminy et al., 1990; Sheng et al., 1990; Van Nguyen et al., 1990). We used double-labeling immunohistochemistry to colocalize CRE activity (i.e., LacZ expression) with orexin or MCH neurons. Chronic administration of morphine caused an ~10-fold induction of CRE activity in orexin cells (Fig. (1A, B) ( $F_{(3,23)} = 49.93$ ; p < 0.05). Precipitation of withdrawal tended to produce a larger induction of CRE activity, although this was not statistically significant. Naltrexone by itself, in morphine-naive animals, produced a much smaller induction of CRE activity in orexin neurons (Fig. 1C,D). Under all three experimental conditions (chronic morphine, morphine withdrawal, or naltrexone alone), there was virtually no induction seen in MCH neurons (Fig. 1A,B). This is in contrast to the selective activation of CRE-LacZ in MCH neurons after starvation (data not shown), indicating that the CRE-LacZ reporter can be activated in both cell types. Interestingly, only a subset of orexin cells showed responses to morphine: 25% of orexin cells were activated after chronic morphine and 33% during withdrawal (Fig. 1B). Moreover, orexin neurons accounted for only approximately one-half of the  $\beta$ -gal-positive neurons observed, indicating that induction of CRE activity also occurs in a population of orexin-negative and MCH-negative neurons.

As a second measure of response to morphine treatments, we studied the induction of c-Fos in wild-type C57BL/6 mice. c-Fos induction has been used as a marker of neuronal activation during opiate withdrawal (Hayward et al., 1990; Georges et al., 2000; Gracy et al., 2001). Whereas chronic morphine treatment, or naltrexone administration to morphine-naive animals, did not induce c-Fos in orexin or MCH cells, precipitation of morphine withdrawal significantly induced c-Fos immunoreactivity in orexin cells, with no response seen in MCH cells (Fig. 1*C*,*D*) ( $F_{(3,20)} = 15.79$ ; p < 0.05). Levels of the Fos family protein FosB, which is also induced by morphine treatments in some brain regions (Nye and Nestler, 1996), were also analyzed. However, we found very little induction of FosB in the LH, with no induction seen in orexin or MCH cells (data not shown).

#### Morphine withdrawal induces orexin gene expression

To assess the impact of chronic morphine and precipitated withdrawal on orexin gene expression, we used a new line of orexinτ-LacZ reporter mice, where the τ-LacZ cassette was knocked into the endogenous or exin locus. The fusion between  $\beta$ -gal and  $\tau$  in these mice makes it possible to visualize neuronal processes in addition to cell bodies. As an initial test of the reporter, one group of mice was deprived of food for 24 hr, and a control group had normal access to food. Food restriction induced a robust increase in the density of LacZ staining in the LH compared with controls (data not shown). Similar increases in orexin mRNA levels have been demonstrated in the LH with food deprivation (Sakurai et al., 1998; Cai et al., 1999; Lopez et al., 2000), indicating that regulation of the reporter gene follows the normal pattern of orexin expression. We subsequently examined the effect of chronic morphine and morphine withdrawal on different groups of orexin-τ-LacZ mice. No change in LacZ staining was detected in response to chronic morphine or to naltrexone in morphinenaive mice (Fig. 2). However, precipitation of morphine withdrawal caused a 56% increase in LacZ staining ( $F_{(3,19)} = 5.78$ ; p <



**Figure 1.** Chronic morphine (Morph) and withdrawal selectively affect orexin (0X) neurons. Confocal microscopy shows double-label immunohistochemistry for orexin or MCH with  $\beta$ -gal (A) or c-Fos (C). The data were quantified and summarized for CRE-LacZ (B) and for c-Fos (D) (n = 6 per group). Nal, Naltrexone. \*p < 0.05.

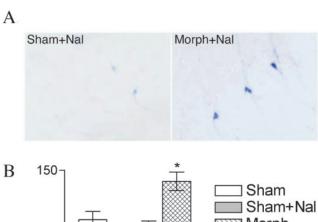
0.01). The increase in LacZ staining during withdrawal was particularly apparent in cellular processes, which demonstrates the utility of the  $\tau$ -LacZ reporter.

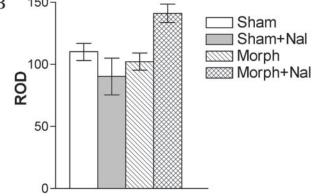
## Morphine-responsive or exin neurons express the $\mu$ -opioid receptor

To investigate the possibility of a direct effect of morphine on orexin cells, we used double-labeling immunofluorescence to examine the presence of MORs on orexin cells in the LH. Examination by conventional and confocal microscopy revealed that ~50% of orexin cells display high levels of MOR immunoreactivity. Because only a subset of orexin cells respond with CRELacZ or c-Fos activation, we performed confocal analysis on triple-immunostained LH sections from animals during precipitated withdrawal. This analysis confirmed that all orexin cells that respond with CRE-LacZ or c-Fos also express MORs, suggesting a direct mode of action. In addition, nonorexin cells that express CRE-LacZ or c-Fos also express MORs (Fig. 3).

# Orexin knock-out mice display attenuated morphine withdrawal

The above molecular studies indicated that orexin neurons and the orexin gene are regulated by morphine treatment. To investigate the possible behavioral significance of this regulation, we studied morphine withdrawal in *orexin* knock-out mice. Precipitation of withdrawal in wild-type littermates induced classic behavioral signs of withdrawal (Fig. 4). Several but not all of these signs were markedly attenuated in the *orexin* knock-outs, and the overall withdrawal score was significantly decreased (wild-type





**Figure 2.** Withdrawal induces orexin gene expression in orexin- $\tau$ -LacZ mice. A, A representative image showing  $\beta$ -gal staining in the LH after sham plus naltrexone (Nal; left) or morphine (Morph) plus naltrexone (right). B, Mean optical densities are indicated (n=5 per group). ROD, Relative optical density.

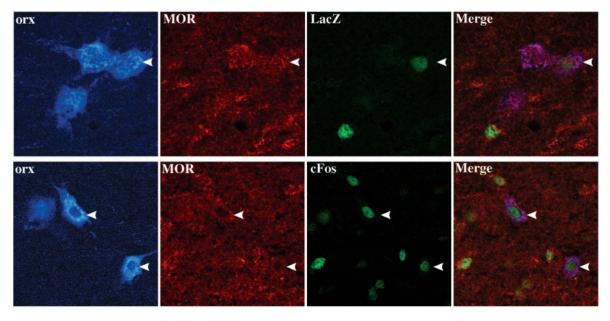


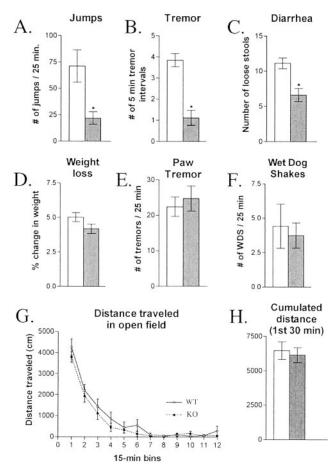
Figure 3. Triple immunostaining showing orexin (orx) cells expressing MOR and LacZ (top) or c-Fos (bottom) as responses to morphine withdrawal. Arrowheads indicate triple positive staining shown on the far right (Merge). Also present are nonorexin cells showing LacZ or c-Fos.

mice, 99.44  $\pm$  15.76; knock-out mice, 42.02  $\pm$  5.38; p < 0.05; n = 8 per group). In contrast, *orexin* mutant mice showed normal open-field locomotor activity (Fig. 4G,H) and stereotypy (data not shown), and no other general behavioral abnormalities were observed.

#### Discussion

Previous ICSS and self-administration studies support the involvement of the LH in behavioral responses to morphine and other opiates (Bozarth et al., 1980; Gallistel et al., 1981). Regions of LH that support morphine self-administration correspond to optimal ICSS sites, which are located in the MFB, a bundle of fibers that connects mesencephalic regions (e.g., the ventral tegmental area) with forebrain structures (e.g., the nucleus accumbens). Although the fibers are known to be important for ICSS, cells intrinsic to the LH also seem to be involved (Velley et al., 1983). The studies reported here are the first to directly implicate orexin neurons in morphine action. Interestingly, endogenous opioids are needed for orexin-induced feeding (Clegg et al., 2002), suggesting multiple points of interaction between orexin and opioids.

Using CRE-LacZ reporter mice, we demonstrate induction of CRE-mediated transcription in the LH by chronic morphine administration, with no additional activation observed during precipitation of withdrawal. A significant fraction of this induction occurs in orexin neurons, whereas virtually no induction is seen in MCH neurons. The fact that CRE activity is seen in the chronic morphine-treated state, without induction of withdrawal, suggests that morphine exposure is sufficient to induce molecular changes that may reflect neural plasticity in response to drug exposure. We also examined induction of c-Fos by chronic morphine and precipitated withdrawal. In contrast to CRE activity, c-Fos was not induced by chronic morphine but was robustly induced during withdrawal. This response, like the induction of CRE activity, was seen in orexin cells but not in MCH cells. These findings suggest that morphine withdrawal produces acute activation of orexin cells. In addition to these changes in transcription factors, morphine withdrawal also induced the orexin gene



**Figure 4.** Morphine withdrawal is attenuated in orexin knock-out mice. *Orexin* knock-outs (gray bars) are compared with their wild-type littermates (white bars) for jumps (A), tremor (B), diarrhea (C), weight loss (D), paw tremor (E), and wet dog shakes (WDS; F) (n=8; \*p<0.01). Open-field locomotor time course (G) and cumulative distance for the first 30 min in a separate set of animals (H) are also shown. ACSF, Artificial CSF.

itself, as seen in orexin- $\tau$ -LacZ reporter mice. Together, these studies demonstrate the regulation of orexin cells by chronic morphine and morphine withdrawal. However, it is important to note that orexin neurons account for only a portion of cells in the LH that show induction of CRE activity and c-Fos in response to chronic morphine or morphine withdrawal. Identification of the other cell types will help to delineate the heterogeneous neuronal populations found in the LH.

The attenuation of morphine withdrawal seen in *orexin* knock-out mice implies that the orexin system serves an important functional role in modulating responses to morphine. The broad projections of orexin neurons and their potential interactions with dopaminergic systems (Nakamura et al., 2000), adrenergic systems (Hagan et al., 1999; Horvath et al., 1999; Bourgin et al., 2000), and the hypothalamic–pituitary–adrenal axis (Kuru et al., 2000) suggest that multiple modes of action may be implicated in orexin modulation of morphine withdrawal. Although *orexin* knock-out mice also display narcolepsy and cataplexy, we did not observe sleep attacks during the 25 min testing period, and locomotor activity was normal. This is expected, because we performed these studies during the light phase, when sleep attacks are infrequent (Chemelli et al., 1999).

Although MORs have been detected previously in the LH (Mansour et al., 1987; Delfs et al., 1994; Mansour et al., 1994; Briski and Sylvester, 2001), the cellular identity of these neurons was unknown. We show in this study that a subset (~50%) of orexin neurons expresses MORs, and that most of these cells respond to morphine by induction of CRE activity or c-Fos. The data also suggest, not surprisingly, that orexin cells are not a homogeneous population. Indeed, it has been demonstrated that amphetamine and antipsychotic drugs induce c-Fos in subsets of orexin neurons (Fadel et al., 2002). It is notable that all cells, both orexin and nonorexin, that show molecular responses to morphine express MORs.

The induction of CRE activity and c-Fos in orexin cells that express the MOR suggests that regulation of the cells by chronic morphine and morphine withdrawal is mediated via direct actions of morphine on these cells. Chronic morphine upregulates the cAMP pathway in many regions of the central and peripheral nervous systems, where it has been shown to be an important mechanism underlying opiate dependence and withdrawal (Nestler, 2001). The expression of MORs on orexin neurons and the activation of CRE-mediated transcription in these cells after chronic morphine treatment suggest that a similar upregulation of the cAMP pathway may occur in these LH neurons. Although additional work is needed to identify the target genes of the cAMP cascade in orexin neurons, the *orexin* gene itself does not appear to be one of them, because chronic morphine did not increase orexin gene expression in the orexin- $\tau$ -LacZ reporter mice. It is also notable that orexin cells express the endogenous opioid peptide dynorphin (Chou et al., 2001); it remains to be determined whether this colocalization has functional implications for mechanisms of morphine withdrawal.

The data presented in this study begin to define, on a molecular and cellular level, the relationship between the LH and drug addiction. Multiple lines of evidence implicate orexin neurons in behavioral responses to morphine. Specifically, induction of c-Fos suggests that orexin neurons are activated during precipitated morphine withdrawal, with the induction of *orexin* gene expression possibly representing a response to restore orexin levels after a period of increased neural activity. The attenuated withdrawal syndrome seen in *orexin* knock-out mice suggests that such regulation of orexin neurons may be an important con-

tributor to morphine physical dependence and to the expression of withdrawal. Additional studies are now needed to assess a role for the orexin system in other behavioral actions of morphine, in particular those actions related to ICSS and self-administration.

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